	RF Errors Corrected by the STIC stems Branch CRF Processing Date: 12 17/2000
Serial N	umber: 09/492,029 Edited by: Verified by: Werlfied by: We
	Changed a file from non-Asoli to New Changed the margins in cases where the sequence text was "wrapped" down to the pext line. IVE Application Data section, specifically:
	Edited a format error in the Current Application Data section, specifically: DEC 2 1 2000 /
	Edited the Current Application Data section with the actual current number. The number inputted by the applicant was the prior application data; or other
<u> </u>	Added the mandatory heading and subheadings for "Current Application Data".
	Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.
	Changed the spelling of a mandatory field (the headings of subheadings), specifically:
	Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were:
	Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited:
	Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.
	Inserted colons after headings/subheadings. Headings edited included:
	Deleted extra, invalid, headings used by an applicant, specifically:
	Deleted: non-ASCII "garbage" at the beginning/end of files; secretary initials/filename at end of file; page numbers throughout text; other invalid text, such as
	Inserted mandatory headings, specifically:
	Corrected an obvious error in the response, specifically:
	Edited identifiers where upper case is used but lower case is required, or vice versa.
	Corrected an error in the Number of Sequences field, specifically:
	A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.
	Deleted ending stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a Patentin bug). Sequences corrected:
回	Other: Seg 2 - corrected (222) response
	to the applicant in the first Office
	A second in the lift Utilico

*Examiner: The above corrections must be communicated to the applicant in the first Office 3/1/95

Action. DO NOT send a copy of this form.

DATE: 12/18/2000 RAW SEQUENCE LISTING TIME: 15:05:02 PATENT APPLICATION: US/09/492,029

Input Set : A:\Pto.amc

Output Set: N:\CRF3\12182000\I492029.raw

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3 <110> APPLICANT: Zuker, Charles S.
        Adler, Jon Elliot
        Lindemeier, Juergen
        The Regents of the University of California
8 <120> TITLE OF INVENTION: Assays for Sensory Modulators Using a Sensory Cell
        Specific G-Protein Beta Subunit
11 <130> FILE REFERENCE: 02307E-092710US
13 <140> CURRENT APPLICATION NUMBER: US 09/492,029
14 <141> CURRENT FILING DATE: 2000-01-26
16 <150> PRIOR APPLICATION NUMBER: US 60/117,404
17 <151> PRIOR FILING DATE: 19 (5) 1-27
19 <160> NUMBER OF SEQ ID NOS: 5
21 <170> SOFTWARE: Patentin Ver
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24 <211> LENGTH: 156
25 <212> TYPE: DNA
26 <213> ORGANISM: Rattus sp.
29 <223> OTHER INFORMATION: rat tongue circumvallate papillae taste receptor
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 52 cecttgacet gtgaace atg ggg gag atg gag cag etg aag cag gag geg
                      Met Gly Glu Met Glu Gln Leu Lys Gin Glu Ala
 53
                                                             10
                                         5
 56 gag cag etc aag aag cag att get gat gee agg aaa gee tgt geg gae
                         1
                                                                        1.58
 57 Glu Gln Leu Lys Lys Gln Ile Ala Asp Ala Arg Lys Ala Cys Ala Asp
                                    20
 60 atc act ctg gct gag ctt gtg tct ggc ctg gag gtg gtg gga cga gtc
                 15
                                                                        206
 61 Ile Thr Leu Ala Glu Leu Val Ser Gly Leu Glu Val Val Gly Arg Val
                                 35
 64 cag atg cgg aca cgg agg acg tta agg gga cac ctg gct aag atc tat
             30
 65 Gln Met Arg Thr Arg Arg Thr Leu Arg Gly His Leu Ala Lys Ile Tyr
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DATE: 12/18/2000 TIME: 15:05:02 RAW SEQUENCE LISTING PATENT APPLICATION: US/09/492,029

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Output Set: N:\CRF3\12182000\I492029.raw

Input Set: A:\f20.dms Output Set: N:\CRF3\12182000\1492029.raw
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68 gcc atg cac tgg gcc act gac tct aag ctg cta gta agt goo act gac tct aag ctg cta gta agt goo act gcc atg cac atg
70 60 65 72 gat ggg aag ctg atc gtg tgg gac act tac acc acc aat aag gtg cat 350 72 gat ggg aag ctg atc gtg tgg gac act tac acc acc aat aag gtg cat 350 72 gat ggg aag ctg atc gtg tgg gac act tac acc acc aat aag gtg cat 350
20 3 an Clu 13/2 Lett 11/2 / 24 "
73 ASP GIV BIS 80 74 80 76 gct atc ccg ctg cgt tcc tcc ttg gtc atg acc tgt qcc tat gca cca 398 76 gct atc ccg ctg cgt tcc tcc ttg gtc atg acc tgt qcc tat gca cca 398
and also the project and see and see
go go ato too too acc
80 tca ggg aac ttc gtg gca tgt ggg ggg sel Asp Asp Met Cys Ser Ile
of the Cly Ach Pile var the -1
an III)
84 tac ago etc aga tec egg gay ago Val Lys Val Ser Arg Glu Leu
ne 195
88 tog gct cac aca ggt tat ctc tcc tgt tgc cgc ttc ccg 42 Asp Asp Asp Asp 89 Ser Ala His Thr Gly Tyr Leu Ser Cys Cys Arg Phe Leu Asp Asp Asp 155
90 140 145 150 150 90 140 145 92 aac att gtg act age tet ggg gac acc acg tgt gcc ttg tgg gac att 590 92 aac att gtg act age tet ggg gac acc acg tgt gcc ttg tgg gac att 590 170 170 170 170 170 170 170 170 170 17
92 aac att gtg act agc tct ggg gac acc acg tgt gcc tcg 93 Asp Ile 92 aac att gtg act agc tct ggg gac acc acg tgt gcc tcg 93 Asp Ile 170 160 165 170 160 1638
94 160 160 165 96 gag acg ggg cag cag aca gtg ttc gtg gga cac act ggt gac tgc 638 96 gag acg ggg cag cag aca gtg ttc gtg gga cac act ggt gac tgc 638
98 175 100 atg agc ctg gct gtg tcc cca gac tac aaa ctc ttc atc tcg gga gct 686 101 Met Ser Leu Ala Val Ser Pro Asp Tyr Lys Leu Phe Ile Ser Gly Ala 101 Met Ser Leu Ala Val Ser Pro 195 195 180 180 180 180 180 180 180 180 195 180 180 180 180 180 180 180 180 180 180
101 Met Sel No. 195 102 190 190 104 tgt gat gcc agc gcc aag ctc tgg gat gtg agg gaa ggg acc tgt cgc 734 104 tgt gat gcc agc gcc aag ctc tgg gat gtg agg gaa ggg acc tgt cgc 734 105 tgg gat gtg agg gaa ggg acc tgt cgc 734
108 cag act ttc act ggc cac yay tee 108 cag act ttc act ggc cac yay tee 108 cag act ttc act ggc cac yay
20 000 - 100 CO
110 220 112 ccc aat ggg gag gcc atc tgc act ggc tca gat gat gat gat gat gat gat gat gat ga
113 Pro Ash Gly Gli Ala 240 245 tog tog gag 878
240 240 240 240 240 245 246 246 247 248 248 248 248 248 248 248 248 248 248
114 116 ctc ttt gac ctg agg gca gac cag gaa ctg aca gct tas ser His Glu 117 Leu Phe Asp Leu Arg Ala Asp Gln Glu Leu Thr Ala Tyr Ser His Glu 117 Leu Phe Asp Leu Arg Ala Asp Gln Glu Leu Thr Ala Tyr Ser His Glu 118 265
118 and atc atc tgt qgc atc acg tcc gta gcc ttc ttc ttc tcu Ser Gly Arg
122 270 124 ctg ctc ttt gct ggc tat gat gac ttc aac tgc adt gtc 393 125 Leu Leu Phe Ala Gly Tyr Asp Asp Phe Asn Cys Asn Val Trp Asp Ser 125 Leu Leu Phe Ala Gly Tyr Asp Asp Phe Asn Cys Asn Val Trp Asp Ser 290 290
128 ctg aag tgt gag cgt gta ggc gtt Ceu Ser Gly His Asp Asn Arg Val
129 Led by 673 and 305 310 130 300 300 305 310 132 agt tgc ctg ggg gtc aca gct gae ggc atg gct gtg gcc act gga tcc 1070
132 agt tgc ctg ggg gtc aca got 3 33

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TENTO IT TO THE

DATE: 12/18/2000 RAW SEQUENCE LISTING PATENT APPLICATION: US/09/492,029 TIME: 15:05:02

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320 320 320 320 111 136 tgg gac age tte etc aag ate tgg gac tgaggagget ggaggagggg 111 136 tgg gac age the lev lys lie Trp Asn	
	7
138 thoughauge atgaagete teagetgact cetatgeet georgetugga ttataggagt 123	7
142 ctataccety gygeadagag gteacaaggg caaagaactg ccccattee technical 144 gtgcctttgg gagtagcagg gteacaaggg caaagaactg ccccattee etcccacce 135 146 ctcctcteea cagtecteat agetteteec tteataaaca agaacagace gagcgeteag 141 146 ctcctcteea cagtecteat agetgett gteeggeetg ggaatggeea statteetge 144 146 147 147 147 147 147 147 147 147 147 147	.7
144 gtqcctttqq quqtaquit agettetece tteataaaca aqaacayatt eteberation 146 cteeteteca cagteetecat agettetece tteataaaca aqaacayatt gtacagetta gaatageca gaatageca gaatageta 141 148 tagatgaca etgggetaca ageagegttt gteeggeetg ggaatageca gaatageta 141 148 tagatgacat etgggetaca teetagtate etggeteect eecagegact ttettetetge 141 151	17
146 ctecteted tageteace ageagegttt gteeggeetg ggaatggeta gugggt 148 tagatgaete etgggetace ageagegttt gteeggeete eecagegaet tretttetge 147 150 eecatgaeta taggtgteac teetagtate etggeteet tegg 152 eecatgaeta taggtgteac teetagtate etggeteet tyg	30
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153 ZOLON TYPE: PRT	
150 -213 ORGANISM: RATTUS SP.	
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162 I. 5 10 164 Glu Ala Asp Ala Arq Lys Ala Cys Ala Asp Ile Thr Leu Ala Glu 30 25 30 Thr Arg	
164 Gln Tle Ala Asp Ala Arq Lys Ala Cys Ala Asp 110	
165 20 25 Val Gln Met Arg Thr Arg	
164 GHI THE MAG 20 165 20 167 Leu Val Ser Gly Leu Glu Val Val Gly Arg Val Gln Met Arg Thr Arg 167 Leu Val Ser Gly Leu Glu Val Val Gly Arg Val Gln Met Arg Thr Arg 168 GH THE MAG 25	
168 35 40 Ala Lys Ile Tyr Ala Met His Trp Ala 170 Arg Thr Leu Arg Gly His Leu Ala Lys Ile Tyr Ala Met His Trp Ala	
170 Arg Thr Leu Arg Gly His Leu Ala Lys IIc 17	
171 50 Ser Gln Asp Gly Lys Leu Tle	
170 Arg fill Led the 55 55 60 171 50 171 Fill Leu Val Ser Ala Ser Gln Asp Gly Lys Leu Ile 173 Thr Asp Ser Lys Leu Leu Val Ser Ala Ser Gln Asp Gly Lys Leu Ile 80 75 75 75 75 75 75 75 75 75 75 75 75 75	
173 THE ASP SET 175 70 174 65 176 Val Trp Asp Thr Tyr Thr Ash Lys Val His Ala Ile Pro Leu Arg 95 90 90	
176 val Trp Asp Thr Tyr Thr Thr Ash Tys 90	
177 85 The Gue Ala Tyr Ala Pro Ser Gly Ash Phe Vai	
176 Val Ttp 135 85 177 85 85 110 105 110 110 105 110 Ser Ser Trp Val Met Thr Cys Ala Tyr Ala Pro Ser Gly Asn Phe Val	
180 100 Acr Met Cys Ser Ile Tyr Ser Leu Lys Ser	
180 100 105 125 126 127 Ser Leu Lys Ser 182 Ala Cys Gly Gly Leu Asp Asn Met Cys Ser Ile Tyr Ser Leu Lys Ser 125 120	
182 Ard Cys Gry Say 120 183 1.15 185 Arg Glu Gly Asn Val Lys Val Ser Arg Glu Leu Ser Ala His Thr Gly 185 Arg Glu Gly Asn Val Lys Val Mbr Ser	
185 Arg Glu Gly Asn var Lys var 135	
186 130 and Cure Arg Phe Leu Asp Asp Asn Asn Ile Val Thi. Ser	
185 Ary Gra Gay 135 186 130 188 Tyr Leu Ser Cys Cys Arg Phe Leu Asp Asp Asn Asn Ile Val Thr Ser 160 150 150 150 160	
189 145 The Obs. Over Ala Leu Trp Asp Ile Glu Thr Gly 1175	
188 Tyl Hed Set 37 150 155 189 145 150 150 155 191 Ser Gly Asp Thr Thr Cys Ala Leu Trp Asp Ile Glu Thr Gly Gln Gln 175 170 170 170 170 170 170 170 170 170 170	
191 Ser Gry Asp 165 165 165 192 194 Lys Thr Val Phe Val Gly His Thr Gly Asp Cys Met Ser Leu Ala Val 194 Lys Thr Val Phe Val Gly His Thr Gly Asp Cys Met Ser Leu Ala Val	
194 Lys Thr Val. Pile Val. 61, 185	
195 180 191 Phe Ile Ser Gly Ala Cys Asp Ala Ser Ala	
185 195 197 180 180 185 197 Ser Pro Asp Tyr Lys Leu Phe Ile Ser Gly Ala Cys Asp Ala Ser Ala 200 200 205 207 207 208 208 209 200 200 207 207 208 208 208 208 208 208 208 208 208 208	
198 195 200 Lys Leu Trp Asp Val Arg Glu Gly Thr Cys Arg Gln Thr Phe Thr Gly 215 220	
200 Lys Leu Trp Asp van 220 220 220	
200 Lys hell Try 1215 201 210 203 His Glu Ser Asp Ile Asn Ala Ile Cys Phe Phe Pro Asn Gly Glu Ala 240 230 230 230 230 240	
203 His Giu Sei Asp 1230 235	
200 Arg Leu Phe Asp Leu Arg 200 Ile Cys Thr Gly Ser Asp Asp Ala Ser Cys Arg Leu Phe Asp Leu Arg 255 265 Ile Cys Thr Gly Ser Asp Asp Ala Ser Cys Arg Leu Phe Asp Leu Arg 255	
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DEC 2 1 2000

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DATE: 12/18/2000 TIME: 15:05:02 RAW SEQUENCE LISTING PATENT APPLICATION: US/09/492,029

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Output Set: N:\CRF3\12182000\1492029.raw

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210 212 The Thr Ser Val Ala Phe Ser Leu Ser Gly Arg Leu Leu Phe Ala Gly 213 275
215 Tyr Asp Asp Phe Asn Cys Asn Val Trp Asp Ser Leu Lys Cys Glu Arg
218 Val Gly Val Leu Ser Gly His Asp Asn Arg Val Ser Cys Leu Gly Val
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221 Thr Ala Asp Gly Met Ala Val Ala Thr Gly Ser Trp Asp Ser Phe Leu
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241 Met Gly Glu Met Clu Gln Leu Arg Gln Glu Ala Glu Gln Leu
242 1 5 10
   244 aag aag cag att gca gat gcc agg aaa gcc tgt gct gac gtt act ctg
                                                                                             96
   245 Lys Lys Gln Ile Ala Asp Ala Arg Lys Ala Cys Ala Asp Val Thr Leu
246 15
   240 13 20 220 248 gca gag ctg gtg tct ggc cta gag gtg gtg gga cga gtc cag atg cgg 248 gca gag ctg gtg tct ggc cta gag gtg gtg gtg aga cga gtc cag atg cgg 249 Ala Glu Leu Val Ser Gly Leu Glu Val Val Gly Arg Val Gln Met Arg 240 35 40
                                                                                             144
   252 acg cgg cgg acg tta agg qga cac ctg gcc aag att tac gcc atg cac
                                                                                              192
    252 and any any and the Leu Arg Gly His Leu Ala Lys Ile Tyr Ala Met His
55 60
    256 tgg gcc act gat tot aag ctg ctg gta agt gcc teg caa gat ggg aag
    257 Trp Ala Thr Asp Ser Lys Leu Leu Val Ser Ala Ser Gln Asp Gly Lys 75 65
    260 ctg atc gtg tgg gac age tac acc acc aac aag gtg cac gcc atc cca
     261 Leu Ile Val Trp Asp Ser Tyr Thr Thr Asn Lys Val His Ala Ile Pro
     264 ctg cgc tcc tcc tgg gtc atg acc tgt gcc tat gcc cca tca ggg aac
                                                                                               336
     204 CLG CGC LCC LGG GGG aLG AGG CGC LGA GGG CGA LGA GGG CAC
265 Leu Arg Ser Ser Trp Val Met Thr Cys Ala Tyr Ala Pro Ser Gly Asn
266 95 100 100 105 105
     268 ttt gtg gca tgt ggg ggg ctg gac aac atg tgt tee atc tac aac etc
                                                                                               384
     200 LLL 9L9 904 L9C 999 999 CCG 980 and acg cyc lee acc lac lac acc lee 269 Phe Val Ala Cys Gly Gly Leu Asp Asn Met Cys Ser Ile Tyr Asn Leu 120 120
     272 aaa too ogt gag ggo aat gto aag gto ago ogg gag ott tot got cao
      272 dad tee ege gag ggs date goo as and Ser Arg Glu Leu Ser Ala His
273 Lys Ser Arg Glu Gly Asn Val Lys Val Ser Arg Glu Leu Ser Ala His
274 130
      276 aca ggt tat etc tec tgc tgc egc ttc etg gat gac aac aal att gtg
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RAW SEQUENCE LISTING DATE: 12/18/2000 PATENT APPLICATION: US/09/492,029 TIME: 15:05:02

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at the mar ten ser Cys Cys Arg Phe Leu Asp Asp 155
277 Thr Gly Tyr Leu Str 77 150 278 150 278 145 280 acc acc acc acc acc acc tgt gcc ttg tgg qac att gag act ggg 528 280 acc acc acc acc acc acc tgt gcc ttg tgg qac att gag act ggg 528 280 acc acc acc acc acc tgt gcc ttg tgg qac att gag act ggg 528 280 acc acc acc acc acc acc acc acc acc ac
278 and tog gag acc acg tgt gcc ttg tgt gag Agn lle Glu Thr Gly
278 145 280 acc age teg gag gac acc acg tegt gee teg teg gac att gag 280 acc age teg gag gac acc acg tegt gee teg teg gac att gag 281 Thr Ser Ser Gly Asp Thr Thr C7s Ala Leu Trp Asp Ile Glu Thr Gly 281 Thr Ser Ser Gly Asp Thr Thr C7s Ala Leu Trp Asp Ile Glu Thr Gly 170 165 170 170 170 170 170 170 170 170 170 170
281 Thr Ser Ser Gry Hor 165 282 160 284 cag cag aag act gta ttt gtg gga cac acg ggt gac tgc atg agc ctg 284 cag cag aag act gta ttt gtg gga cac acg ggt gac tgc atg agc ctg 284 cag cag aag act gta ttt gtg gga cac acg ggt gac tgc atg agc ctg 287 Thr Ser Ser Gry Hor Thr Glog acc acg ggt gac tgc atg agc ctg 288 Cag cag aag act gta ttt gtg gga cac acg ggt gac tgc atg agc ctg 289 Thr Ser Ser Gry Hor Thr Glog acc acg ggt gac tgc atg agc ctg 280 Thr Ser Ser Gry Hor Thr Glog acc acg ggt gac tgc atg agc ctg 280 Thr Ser Ser Gry Hor Thr Glog acc acg ggt gac tgc atg agc ctg 281 Thr Ser Ser Gry Hor Thr Glog acc acg ggt gac tgc atg agc ctg 282 Thr Glog acc acg ggt gac tgc atg agc ctg 283 Thr Glog acc acg ggt gac tgc atg agc ctg 284 Cag cag aag act gta ttt gtg gga cac acg ggt gac tgc atg agc ctg 285 Thr Glog Asp Cys Net Ser Leu 286 Thr Glog acc acg ggt gac tgc atg agc ctg 286 Thr Glog acc acg ggt gac tgc atg agc ctg 287 Thr Glog acc acg ggt gac tgc atg agc ctg 288 Thr Glog acc acg ggt gac tgc atg acc acg ggt gac tgc atg acc acg ggt gac tgc atg agc ctg
282 160 act gta ttt gtg gga cac acg ggt gue chan cys Net ser Leu
282 160 284 cag cag aag act gta ttt gtg gga cac acg ggt gac tyc acg 285 Gln Gln Lys Thr Val Phe Val Gly His Thr Gly Asp Cys Met Ser Leu 285 Gln Gln Lys Thr Val Phe Val Gly His Thr Gly Asp Cys Met Ser Leu 190 185 186 624
285 Gln Gla Lys III 180 180 agg gcc tqt gat gcc 624
285 Gln Gln Lys Thi VI 180 286 175 288 gct gtg tct cct gac ttc aat ctc ttc att tcg ggg gcc tgt gat gcc 624 288 gct gtg tct cct gac ttc aat ctc ttc att tcg ggg gcc tgt gat gcc 624 288 gct gtg tct cct gac ttc aat ctc ttc att tcg ggg gcc tgt gat gcc 624 285 gct gtg tct cct gac ttc aat ctc ttc att tcg ggg gcc tgt gat gcc 624 285 gct gtg tct cct gac ttc aat ctc ttc att tcg ggg gcc tgt gat gcc 624 285 gct gtg tct cct gac ttc aat ctc ttc att tcg ggg gcc tgt gat gcc 624 286 gct gtg tct cct gac ttc aat ctc ttc att tcg ggg gcc tgt gat gcc 624 286 gct gtg tct cct gac ttc aat ctc ttc att tcg ggg gcc tgt gat gcc 624 286 gct gtg tct cct gac ttc aat ctc ttc att tcg ggg gcc tgt gat gcc 624 286 gct gtg tct cct gac ttc aat ctc ttc att tcg ggg gcc tgt gat gcc 624
286 175 288 gct gtg tct cct gac ttc aat ctc ttc att tcg ggg gcc tgc gcd Ala Cys Asp Ala 289 Ala Val Ser Pro Asp Phe Asn Leu Phe Ile Ser Gly Ala Cys Asp Ala 205 200 200 200 201 200 200 201 200 200 200
289 Ala Val Ser F10 Kap 195 200 290 195 292 agt gcc aag ctc tqg gat gtg cga gag qgg acc tqc cgt cag act ttc 672 292 agt gcc aag ctc tqg gat gtg cga gag qgg acc tqc cgt cag act ttc 672 292 agt gcc aag ctc tqg gat gtg cga gag qgg acc tqc cgt cag act ttc 672 292 agt gcc aag ctc tqg gat gtg cga gag qgg acc tqc cgt cag act ttc 672 292 agt gcc aag ctc tqg gat gtg cga gag ggg acc tqc cgt cag act ttc 672 292 agt gcc aag ctc tqg gat gtg cga gag ggg acc tqc cgt cag act ttc 672 292 agt gcc aag ctc tqg gat gtg cga gag ggg acc tqc cgt cag act ttc 672 292 agt gcc aag ctc tqg gat gtg cga gag ggg acc tqc cgt cag act ttc 672 292 agt gcc aag ctc tqg gat gtg cga gag ggg acc tqc cgt cag act ttc 672
290 agt gcc agg ctc tgg gat gtg cga gag ggg agg agg ggg agg Glu Thr Phe
290 292 agt gcc aag ctc tgg gat gtg cga gag ggg acc tgc cgc ctg ctg ctg cgc 292 agt gcc aag ctc tgg gat gtg cga gag ggg acc tgc cgc cag ctg ctg ctg cag cag cgc agt gga ggg acc tgc cgc cag ctg ctg ctg ctg ctg ctg ctg ctg ctg ct
293 Ser Ala Lys her Try Nor 215 294 296 act ggc cac gag tog gac atc aac gcc atc tgt ttc tcc coc aat gga 720 296 act ggc cac gag tog gac atc aac gcc atc tgt ttc tcc coc aat gga 720 296 act ggc cac gag tog gac atc aac gcc atc tgt ttc tcc coc aat gga 720 298 act ggc cac gag tog gac atc acc tgt ttc tcc coc aat gga 720 298 act ggc cac gag tog gac atc acc tgt ttc tcc coc aat gga 720
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294 296 act ggc cac gag tog gac ate aac gcc ate tgt tte tee to the act ggc act gag tog gac ate aac gcc ate tgt tte tee tot act ggc act gag tog gac ate act gcc ate tgt tte tee tot gag act gact g
297 Thr Gly His Git 352 230 230 230 tcc tgc cgc ttg ttt qac 768
297 Thr Gly His Git Ser Nov 230 298 225 300 gag qcc atc tgc acg ggc tcg gat gac gct tcc tgc cqc ttg ttt qac 768 300 gag qcc atc tgc acg ggc tcg gat gac gct tcc tgc cqc ttg ttt qac 768 300 gag qcc atc tgc acg ggc tcg gat gac gct tcc tgc cqc ttg ttt qac 768
298 225 300 gag qcc atc tgc acg ggc tcg gat gac gct tcc tgc tqc tcg tqc tcg 301 Glu Ala Ile Cys Thr Gly Ser Asp Asp Ala Ser Cys Arg Leu Phe Asp 250 245 245 250 267 268 268 268 268 268 268 268 268 268 268
301 Glu Ala IIe Cys Ini 245 302 240 304 ctg cgg gca gac cag gag ctg atc tgc ttc tcc cac gag agc atc atc 816 304 ctg cgg gca gac cag gag ctg atc tgc ttc tcc cac gag agc atc atc 816 304 ctg cgg gca gac cag gag ctg atc tgc ttc tcc cac gag agc atc atc 816 305 245 307 81a Asp Glo Glu Leu IIe Cys Phe Ser His Glu Ser IIe IIe 270
304 ctg cqq gca gac cag gag ctg atc cys Phe Ser His Glu Ser Ile Ile
302 240 304 ctg cgg gca gac cag gag ctg atc tgc ttc tcc cac gag agc tg atc tgc ttc tcc cac gag agc tg atc tgc ttc tcc cac gag agc tg atc tgc tcc tcc cac gag agc tg atc tle Ile 304 ctg cgg gca gac cag gag ctg atc tgc tcc tcc cac gag agc tgc tle Ile 305 Leu Arg Ala Asp Gin Glu Leu Ile Cys Phe Ser His Glu Ser Ile Ile 270 265 265 266 268 268 268 268 268 268 268 268 268
306 255 260 age the tee etc agt gge ege eta eta tite to
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VERIFICATION SUMMARY DATE: 12/18/2000 TIME: 15:05:03

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Output Set: N:\CRF3\12182000\1492029.raw